



## SEQUENCE LISTING

<110> KISELEV, VSEVOLOD I  
PETR, SVESHNIKOV G

<120> METHODS, KITS, AND COMPOSITIONS FOR THE DEVELOPMENT AND  
USE OF MONOCLONAL ANTIBODIES SPECIFIC TO ANTIGENS  
TRADITIONALLY OF LOW IMMUNOGENICITY

<130> 16631.0001

<140> 10/573,478

<141> 2006-03-24

<150> PCT/RU2004/000373

<151> 2004-09-24

<150> RU 2003128660

<151> 2003-09-25

<160> 22

<170> PatentIn Ver. 3.3

<210> 1

<211> 309

<212> DNA

<213> Human papillomavirus type 16

<220>

<221> CDS

<222> (7)..(303)

<400> 1

gaattc atc atg cat gga gat aca cct aca ttg cat gaa tat atg tta	48
Ile Met His Gly Asp Thr Pro Thr Leu His Glu Tyr Met Leu	
1 5 10	
gat ttg caa cca gag aca act gat ctc tac tgt tat gag caa tta aat	96
Asp Leu Gln Pro Glu Thr Thr Asp Leu Tyr Cys Tyr Glu Gln Leu Asn	
15 20 25 30	
gac agc tca gag gag gag gat gaa ata gat ggt cca gct gga caa gca	144
Asp Ser Ser Glu Glu Glu Asp Glu Ile Asp Gly Pro Ala Gly Gln Ala	
35 40 45	
gaa ccg gac aga gcc cat tac aat att gta acc ttt tgt tgc aag tgt	192
Glu Pro Asp Arg Ala His Tyr Asn Ile Val Thr Phe Cys Cys Lys Cys	
50 55 60	
gac tct acg ctt cgg ttg tgc gta caa agc aca cac gta gac att cgt	240
Asp Ser Thr Leu Arg Leu Cys Val Gln Ser Thr His Val Asp Ile Arg	
65 70 75	
act ttg gaa gac ctg tta atg ggc aca cta gga att gtg tgc ccc atc	288
Thr Leu Glu Asp Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro Ile	
80 85 90	

tgt tct cag aaa cca ggatcc  
 Cys Ser Gln Lys Pro  
 95

309

<210> 2  
 <211> 99  
 <212> PRT  
 <213> Human papillomavirus type 16

<400> 2  
 Ile Met His Gly Asp Thr Pro Thr Leu His Glu Tyr Met Leu Asp Leu  
 1 5 10 15  
 Gln Pro Glu Thr Thr Asp Leu Tyr Cys Tyr Glu Gln Leu Asn Asp Ser  
 20 25 30  
 Ser Glu Glu Glu Asp Glu Ile Asp Gly Pro Ala Gly Gln Ala Glu Pro  
 35 40 45  
 Asp Arg Ala His Tyr Asn Ile Val Thr Phe Cys Cys Lys Cys Asp Ser  
 50 55 60  
 Thr Leu Arg Leu Cys Val Gln Ser Thr His Val Asp Ile Arg Thr Leu  
 65 70 75 80  
 Glu Asp Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro Ile Cys Ser  
 85 90 95  
 Gln Lys Pro

<210> 3  
 <211> 330  
 <212> DNA  
 <213> Human papillomavirus type 18

<220>  
 <221> CDS  
 <222> (7)..(324)

<400> 3  
 gaattc agt atg cat gga cct aag gca aca ttg caa gac att gta ttg 48  
 Ser Met His Gly Pro Lys Ala Thr Leu Gln Asp Ile Val Leu  
 1 5 10  
 cat tta gag ccc caa aat gaa att ccg gtt gac ctt cta tgt cac gag 96  
 His Leu Glu Pro Gln Asn Glu Ile Pro Val Asp Leu Leu Cys His Glu  
 15 20 25 30  
 caa tta agc gac tca gag gaa gaa aac gat gaa ata gat gga gtt aat 144  
 Gln Leu Ser Asp Ser Glu Glu Glu Asn Asp Glu Ile Asp Gly Val Asn  
 35 40 45  
 cat caa cat tta cca gcc cga cga gct gaa cca caa cgt cac aca atg 192  
 His Gln His Leu Pro Ala Arg Arg Ala Glu Pro Gln Arg His Thr Met  
 50 55 60

ttg tgt atg tgt tgt aag tgt gaa gcc aga att gag cta gta gta gaa 240  
 Leu Cys Met Cys Cys Lys Cys Glu Ala Arg Ile Glu Leu Val Val Glu  
           65                          70                          75

agc tca gca gac gac ctt cga gca ttc cag cag ctg ttt ctg aac acc 288  
 Ser Ser Ala Asp Asp Leu Arg Ala Phe Gln Gln Leu Phe Leu Asn Thr  
           80                          85                          90

ctg tcc ttt gtg tgt ccg tgg tgt gca tcc cag cag ggatcc 330  
 Leu Ser Phe Val Cys Pro Trp Cys Ala Ser Gln Gln  
           95                          100                          105

<210> 4

<211> 106

<212> PRT

<213> Human papillomavirus type 18

<400> 4

Ser Met His Gly Pro Lys Ala Thr Leu Gln Asp Ile Val Leu His Leu  
           1                          5                          10                          15

Glu Pro Gln Asn Glu Ile Pro Val Asp Leu Leu Cys His Glu Gln Leu  
                           20                          25                          30

Ser Asp Ser Glu Glu Glu Asn Asp Glu Ile Asp Gly Val Asn His Gln  
           35                          40                          45

His Leu Pro Ala Arg Arg Ala Glu Pro Gln Arg His Thr Met Leu Cys  
           50                          55                          60

Met Cys Cys Lys Cys Glu Ala Arg Ile Glu Leu Val Val Glu Ser Ser  
           65                          70                          75                          80

Ala Asp Asp Leu Arg Ala Phe Gln Gln Leu Phe Leu Asn Thr Leu Ser  
                           85                          90                          95

Phe Val Cys Pro Trp Cys Ala Ser Gln Gln  
           100                          105

<210> 5

<211> 5321

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic nucleotide  
 sequence of recombinant vector pQE30-dnaK

<400> 5

ctcgagaaat cataaaaaat ttatttgctt tgtgagcgga taacaattat aatagattca 60  
 attgtgagcg gataacaatt tcacacagaa ttcattaaag aggagaaatt aactatgaga 120  
 ggatcgcatc accatcacca tcacggatcc gctcgtgcgg tcgggatcga cctcgggacc 180  
 accaactccg tcgtctcggg tctggaaggt ggcgacccgg tcgtcgtcgc caactccgag 240  
 ggctccagga ccaccccgtc aattgtcgcg ttgcgccgca acggtgaggt gctggtcggc 300  
 cagcccccca agaaccaggc agtgaccaac gtcgatcgca ccgtgcgctc ggtcaagcga 360

cacatgggca	gogactgggc	catagagatt	gacggcaaga	aatacacccg	gccggagatc	420
agcgcccgca	ttctgatgaa	gctgaagcgc	gacgccgagg	cctacctcgg	tgaggacatt	480
accgacgcgg	ttatcacgac	gcccgcctac	ttcaatgacg	cccagcgtca	ggccaccaag	540
gacgccggcc	agatcgccgg	cctcaacgtg	ctgcccgatcg	tcaacgagcc	gaccgcggcc	600
gcgctggcct	acggcctcga	caagggcgag	aaggagcagc	gaatcctggt	cttcgacttg	660
ggtggtggca	ctttcgacgt	ttccctgctg	gagatcggcg	aggggtgtgt	tgagggtccgt	720
gccacttcgg	gtgacaacca	cctcggcggc	gacgactggg	accagcgggt	cgtcgatttg	780
ctggtggaca	agttcaaggg	caccagcggc	atcgatctga	ccaaggacaa	gatggcgatg	840
cagcggctgc	gggaagccgc	cgagaaggca	aagatcgagc	tgagttcgag	tcagtcacc	900
tcgatcaacc	tgccctacat	caccgtcgac	gccgacaaga	acccgttggt	cttagacgag	960
cagctgaccc	gcgcggaggt	ccaacggatc	actcaggacc	tgctggaccg	cactcgcaag	1020
ccgttccagt	cggtgatcgc	tgacaccggc	atttcgggtg	cggagatcga	tcacgttggt	1080
ctcgtgggtg	gttcgacccg	gatgcccgcg	gtgaccgatc	tggtcaagga	actcaccggc	1140
ggcaaggaac	ccaacaaggg	cgtcaacccc	gatgaggttg	tcgcgggtgg	agccgctctg	1200
caggccggcg	tcctcaaggg	cgaggtgaaa	gacgttctgc	tgcttgatgt	taccccgctg	1260
agcctgggta	tcgagaccaa	gggcgggggt	atgaccaggc	tcatecgagc	caacaccacg	1320
atccccacca	agcggtcgga	gactttcacc	accgccgacg	acaaccaacc	gtcgggtgac	1380
atccaggctc	atcaggggga	gcgtgagatc	gccgcgcaca	acaagttgct	cgggtccctc	1440
gagctgaccg	gcatcccgcg	ggcgcgcggg	gggattccgc	agatcgaggt	cactttcgac	1500
atcgacgcca	acggcattgt	gcacgtcacc	gccaaggaca	agggcaccgg	caaggagaac	1560
acgatccgaa	tcaggaagg	ctcgggcctg	tccaagggaag	acattgaccg	catgatcaag	1620
gacgccgaag	cgcacgccga	ggaggatcgc	aagcgtcgcg	aggaggccga	tgttcgtaat	1680
caagccgaga	cattgggtcta	ccagacggag	aagttcgtca	aagaacagcg	tgaggccgag	1740
ggtggttcga	aggtacctga	agacacgctg	aacaagggtt	atgccgcggg	ggcgggaagcg	1800
aaggcggcac	ttggcggatc	ggatatttcg	gccatcaagt	cggcgatgga	gaagctgggc	1860
caggagtgcg	aggctctggg	gcaagcgatc	tacgaagcag	ctcaggctgc	gtcacaggcc	1920
actggcgag	cccaccccg	cggcgagccg	ggcgggtgcc	accccggtc	ggctgatgac	1980
gttggtggac	cggagggtgt	cgacgacggc	cgggaaggcca	agtgacggac	gggtcgacct	2040
gcagccaagc	ttaattagct	gagcttggtg	tcctgttgat	agatccagta	atgacctcag	2100
aactccatct	ggatttggtc	agaacgctcg	gttgccgcgc	ggcgtttttt	attggtgaga	2160
atccaagcta	gcttggcgag	attttcagga	gctaagggaag	ctaaaatgga	gaaaaaaatc	2220
actggatata	ccaccgttga	tatatcccaa	tggcatcgta	aagaacattt	tgaggcattt	2280
cagtcagttg	ctcaatgtac	ctataaccag	accgttcagc	tggatattac	ggccttttta	2340
aagaccgtaa	agaaaaataa	gcacaagttt	tatccggcct	ttattcacat	tcttgcccg	2400
ctgatgaatg	ctcatccgga	atttcgatat	gcaatgaaag	acggtgagct	ggtgatattg	2460
tgagtgttc	acccttggtt	caccgttttc	catgagcaaa	ctgaaacgtt	ttcatcgctc	2520
tggagtgaat	accacgacga	tttcggcgag	tttctacaca	tatattcgca	agatgtggcg	2580
tgttacggtg	aaaacctggc	ctatttcctt	aaagggttta	ttgagaatat	gtttttcgtc	2640
tcagccaatc	cctgggtgag	tttcaccagt	tttgatttaa	acgtggccaa	tatggacaac	2700
ttcttcgccc	ccgttttcac	catgggcaaa	tattatacgc	aaggcgacaa	ggtgctgatg	2760
ccgctggcga	ttcaggttca	tcatgccgtt	tgtgatggct	tccatgtcgg	cagaatgctt	2820
aatgaattac	aacagtactg	cgatgagtg	cagggcgggg	cgtaattttt	ttaaggcagt	2880
tattggtgcc	cttaaaccgc	tggggtaatg	actctctagc	ttgaggcatc	aaataaaacg	2940
aaaggctcag	tcgaaagact	gggcctttcg	ttttatctgt	tgtttgctcg	tgaacgctct	3000
cctgagttag	acaaatccgc	cctctagagc	tgccctgcgc	gtttcggtga	tgacggtgaa	3060
aacctctgac	acatgcagct	cccggagacg	gtcacagctt	gtctgtaagc	ggatgccggg	3120
agcagacaag	cccgctcagg	cgcgtcagcg	ggtgttggtg	ggtgtcgggg	cgcagccatg	3180
acccagtcac	gtagcgatag	cggagtgtat	actggcttaa	ctatgcggca	tcagagcaga	3240
ttgtactgag	agtgcaccat	atgcggtgtg	aaataaccgca	cagatgcgta	aggagaaaat	3300
accgcacag	gcgctcttcc	gcttcctcgc	tactgactc	gctgcgctcg	gtcgttcggc	3360
tgccggcgag	ggtatcagct	cactcaaagg	cggtaatac	gttatccaca	gaatcagggg	3420
ataacgcagg	aaagaacatg	tgagcaaaa	gccagcaaaa	ggccaggaac	cgtaaaaagg	3480
ccgctgtgtc	ggcggttttc	cataggctcc	gccccctga	cgagcatcac	aaaaatcgac	3540
gctcaagtca	gaggtggcga	aaccgcagc	gactataaag	ataccaggcg	tttccccctg	3600
gaagctccct	cgtgcctctc	cctgttcga	ccctgcgct	taccggatac	ctgtccgctc	3660
ttctcccttc	gggaagcggt	gcgctttctc	atagctcag	ctgtaggtat	ctcagttcgg	3720
tgtaggctcg	tcgctccaag	ctgggctgtg	tgcacgaacc	ccccgttcag	cccgaccgct	3780
gcgccttctc	cggttaactat	cgtcttgagt	ccaaccgggt	aagacacgac	ttatcgccac	3840

```

tggcagcagc cactggtaac aggattagca gagcgaggta tgtaggcggt gctacagagt 3900
tcttgaagtg gtggcctaac tacggctaca ctagaaggac agtatttggg atctgcgctc 3960
tgctgaagcc agttaccttc ggaaaaagag ttggtagctc ttgatccggc aaacaaacca 4020
ccgctggtag cgggtggtttt tttgtttgca agcagcagat tacgcgcaga aaaaaaggat 4080
ctcaagaaga tcctttgatc ttttctacgg ggtctgacgc tcagtggaac gaaaactcac 4140
gttaagggat tttggtcatt agattatcaa aaaggatctt cacctagatc cttttaaat 4200
aaaaatgaag ttttaaatca atctaaagta tatatgagta aacttggtct gacagttacc 4260
aatgcttaat cagtgaggca cctatctcag cgatctgtct atttcgttca tccatagttg 4320
cctgactccc cgctgtgtag ataactacga tacgggaggg cttaccatct ggccccagt 4380
ctgcaatgat accgcgagac ccacgctcac cggctccaga tttatcagca ataaaccagc 4440
cagccggaag ggccgagcgc agaagtggct ctgcaacttt atccgcctcc atccagtcta 4500
ttaattgttg ccgggaagct agagtaagta gttcgccagt taatagtttg cgcaacgttg 4560
ttgccattgc tacaggcatc gtggtgtcac gctcgtcgtt tggtagggct tcattcagct 4620
ccggttccca acgatcaagg cgagttacat gatcccccat gttgtgcaaa aaagcggtta 4680
gctccttcgg tcctccgatc gttgtcagaa gtaagttggc cgcagtgtta tcactcatgg 4740
ttatggcagc actgcataat tctcttactg tcatgccatc cgtaagatgc ttttctgtga 4800
ctggtagagta ctcaaccaag tcattctgag aatagtgtat gcggcgaccg agttgctctt 4860
gcccgcgctc aatacgggat aataccgcgc cacatagcag aactttaaaa gtgctcatca 4920
ttggaaaacg ttcttcgggg cgaaaactct caaggatctt accgctgttg agatccagtt 4980
cgatgtaacc cactcgtgca cccaactgat cttcagcatc ttttactttc accagcgttt 5040
ctgggtgagc aaaaacagga aggcaaaatg ccgcaaaaaa gggaataagg gcgacacgga 5100
aatgttgaat actcactatc ttcctttttc aatattattg aagcatttat cagggttatt 5160
gtctcatgag cggatacata tttgaatgta tttagaaaaa taaacaaata ggggttccgc 5220
gcacatttcc ccgaaaagtg ccacctgacg tctaagaaac cattattatc atgacattaa 5280
cctataaaaa taggcgtatc acgaggccct ttcgtcttca c 5321

```

```

<210> 6
<211> 12
<212> PRT
<213> Bos taurus

```

```

<400> 6
Lys Lys Arg Pro Lys Pro Gly Gly Gly Trp Asn Thr
  1                      5                      10

```

```

<210> 7
<211> 8
<212> PRT
<213> Bos taurus

```

```

<400> 7
Gln Pro His Gly Gly Gly Trp Gly
  1                      5

```

```

<210> 8
<211> 13
<212> PRT
<213> Bos taurus

```

```

<400> 8
Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Ile Lys
  1                      5                      10

```

<210> 9  
 <211> 17  
 <212> PRT  
 <213> Bos taurus

<400> 9  
 Ile Thr Gln Tyr Gln Arg Glu Ser Gln Ala Tyr Tyr Gln Arg Gly Ala  
       1                  5                  10                  15

Ser

<210> 10  
 <211> 19  
 <212> DNA  
 <213> Human papillomavirus type 16

<400> 10  
 tgacagctca gaggaggag 19

<210> 11  
 <211> 19  
 <212> DNA  
 <213> Human papillomavirus type 16

<400> 11  
 gcacaaccga agcgtagag 19

<210> 12  
 <211> 20  
 <212> DNA  
 <213> Human papillomavirus type 18

<400> 12  
 gcgactcaga ggaagaaaac 20

<210> 13  
 <211> 20  
 <212> DNA  
 <213> Human papillomavirus type 18

<400> 13  
 caaaggacag ggtgttcaga 20

<210> 14  
 <211> 31  
 <212> DNA  
 <213> Human papillomavirus type 18

<400> 14  
 tctaacgaat tcagtatgca tggacctaag g 31

<210> 15  
 <211> 30  
 <212> DNA  
 <213> Human papillomavirus type 18

<400> 15  
 attacaggat ccctgctggg atgcacacca 30

<210> 16  
 <211> 31  
 <212> DNA  
 <213> Human papillomavirus type 16

<400> 16  
 attctcgaat tcatcatgca tggagatata c 31

<210> 17  
 <211> 31  
 <212> DNA  
 <213> Human papillomavirus type 16

<400> 17  
 cttatcggat cctggtttct gagaacagat g 31

<210> 18  
 <211> 130  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic pHE716 and  
 pHE718 terminal sequence

<220>  
 <221> misc\_feature  
 <222> (107)..(108)  
 <223> HSP 16/HSP18 E7 gene insertion site

<400> 18  
 taatacgact cactataggg agaccacaac ggtttccctc tagaaataat tttgtttaac 60  
 tttaagaagg agatatacat atgcatcacc atcaccatca cgaattcgga tcctaattag 120  
 ctgaaagctt 130

<210> 19  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 primer

<400> 19  
 gaagatctat gcatggagat acacctac 28

<210> 20  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
primer

<400> 20  
cgggatcctg gtttctgaga acagatgg 28

<210> 21  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
primer

<400> 21  
gaagatctat gcatggacct aaggcaac 28

<210> 22  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
primer

<400> 22  
cgggatccct gctgggatgc acaccacg 28